

TABLE 3T Genes Specific for Allergy Only 257

Spot	p-value	Description	Accession	Unigene
140	0.025999	yg33d08.s1 Soares infant brain 1NIB cDNA clone IMAGE:34540 3', mRNA sequence /clone=IMAGE:34540 /clone_end=3' /gb=R44968 /gi=824322 /ug=Hs.412527 /len=453	R44968	Hs.412527
377	0.009301	basic helix-loop-helix domain containing, class B, 3 (BHLHB3), mRNA /cds=(135,1583) /gb=NM_030762 /gi=13540520 /ug=Hs.33829 /len=3641	NM_030762	Hs.33829
424	0.025999	zinc-fingers and homeoboxes 1 (ZHX1), mRNA /cds=(25,2646) /gb=NM_007222 /gi=6005959 /ug=Hs.12940 /len=4339	NM_007222	Hs.12940
428	0.04114	golgi complex associated protein 1, 60kDa (GOCAP1), mRNA /cds=(56,1642) /gb=NM_022735 /gi=15826851 /ug=Hs.6831 /len=3598	NM_022735	Hs.6831
725	0.03394	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 24 (DDX24), mRNA /cds=(100,2679) /gb=NM_020414 /gi=14251213 /ug=Hs.155986 /len=2967	NM_020414	Hs.155986
736	0.02782	myosin, heavy polypeptide 11, smooth muscle (MYH11), transcript variant SM2, mRNA /cds=(89,5905) /gb=NM_022844 /gi=13124874 /ug=Hs.78344 /len=6900	NM_022844	Hs.78344
1051	0.049555	HSPC133 protein (HSPC133), mRNA /cds=(83,481) /gb=NM_014168 /gi=7661791 /ug=Hs.273063 /len=963	NM_014168	Hs.273063
1076	0.036213	ribosomal protein S20 (RPS20), mRNA /cds=(128,487) /gb=NM_001023 /gi=14591915 /ug=Hs.8102 /len=539	NM_001023	Hs.8102
1272	0.049555	collagen, type XII, alpha 1 (COL12A1), transcript variant long, mRNA /cds=(114,9305) /gb=NM_004370 /gi=18201922 /ug=Hs.101302 /len=11554	NM_004370	Hs.101302
1412	0.04114	lipoma HMGIC fusion partner (LHFP), mRNA /cds=(357,959) /gb=NM_005780 /gi=5031864 /ug=Hs.93765 /len=2012	NM_005780	Hs.93765
1561	0.043803	translocating chain-associating membrane protein (TRAM), mRNA /cds=(92,1216) /gb=NM_014294 /gi=19923404 /ug=Hs.4147 /len=2722	NM_014294	Hs.4147
1653	0.036213	peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA /cds=(45,542) /gb=NM_021130 /gi=10863926 /ug=Hs.401787 /len=753	NM_021130	Hs.401787
1691	0.013664	zinc finger protein 36, C3H type-like 1 (ZFP36L1), mRNA /cds=(131,1147) /gb=NM_004926 /gi=15812179 /ug=Hs.85155 /len=3022	NM_004926	Hs.85155
1947	0.038612	transforming, acidic coiled-coil containing protein 1 (TACC1), mRNA /cds=(321,2738) /gb=NM_006283 /gi=5454099 /ug=Hs.173159 /len=7758	NM_006283	Hs.173159
2104	0.013664	hPMS3 mRNA, partial cds. /cds=(1,773) /gb=D38435 /gi=600590 /ug=Hs.334451 /len=846	D38435	Hs.334451
2176	0.03394	proteasome (prosome, macropain) 26S subunit, non- ATPase, 12 (PSMD12), mRNA /cds=(44,1414) /gb=NM_002816 /gi=4506220 /ug=Hs.4295 /len=3548	NM_002816	Hs.4295

2270	0.019683	MTB (MTB) mRNA, complete cds /cds=(80,265) /gb=AF348994 /gi=28190031 /ug=Hs.333727 /len=408	AF348994	Hs.333727
2329	0.043803	inhibitor of growth family, member 1-like (ING1L), mRNA /cds=(92,934) /gb=NM_001564 /gi=4504694 /ug=Hs.107153 /len=1078	NM_001564	Hs.107153
2370	0.043803	exosome component Rrp41 (FLJ20591), mRNA /cds=(104,841) /gb=NM_019037 /gi=9506688 /ug=Hs.343589 /len=896	NM_019037	Hs.343589
2635	0.036213	NS1-binding protein (NS1-BP), mRNA /cds=(556,2484) /gb=NM_006469 /gi=24475846 /ug=Hs.197298 /len=4137	NM_006469	Hs.197298
2687	0.043803	v-fos FBJ murine osteosarcoma viral oncogene (FOS), mRNA /cds=(156,1298) /gb=NM_005252 /gi=6552332 /ug=Hs.25647 /len=2084	NM_005252	Hs.25647
2826	0.02782	potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3 (KCNS3), mRNA /cds=(381,1856) /gb=NM_002252 /gi=25952107 /ug=Hs.47584 /len=2344	NM_002252	Hs.47584
2853	0.038612	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 1, 7kDa (NDUFB1), mRNA /cds=(34,210) /gb=NM_004545 /gi=27597086 /ug=Hs.183435 /len=290	NM_004545	Hs.183435
2858	0.043803	hypothetical protein HSPC228 (HSPC228), mRNA /cds=(16,939) /gb=NM_016485 /gi=21361740 /ug=Hs.267288 /len=3273	NM_016485	Hs.267288
2860	0.029748	peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA /cds=(45,542) /gb=NM_021130 /gi=10863926 /ug=Hs.401787 /len=753	NM_021130	Hs.401787
2898	0.029748	zinc finger protein 140 (clone pHZ-39) (ZNF140), mRNA /cds=(273,1646) /gb=NM_003440 /gi=4507990 /ug=Hs.154205 /len=2407	NM_003440	Hs.154205
2901	0.046607	A kinase (PRKA) anchor protein (yotiao) 9 (AKAP9), transcript variant 4, mRNA /cds=(223,5190) /gb=NM_147166 /gi=22538388 /ug=Hs.58103 /len=6058	NM_147166	Hs.58103
2931	0.043803	peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA /cds=(45,542) /gb=NM_021130 /gi=10863926 /ug=Hs.401787 /len=753	NM_021130	Hs.401787
3062	0.036213	myosin, light polypeptide, regulatory, non-sarcomeric (20kD) (MLCB), mRNA /cds=(115,630) /gb=NM_006471 /gi=5453739 /ug=Hs.180224 /len=944	NM_006471	Hs.180224
3082	0.017049	speckle-type POZ protein (SPOP), mRNA /cds=(158,1282) /gb=NM_003563 /gi=4507182 /ug=Hs.129951 /len=1642	NM_003563	Hs.129951
3117	0.046607	mRNA for Sec24 protein (Sec24A isoform), partial /cds=(1,3237) /gb=AJ131244 /gi=3947687 /ug=Hs.211612 /len=5967	AJ131244	Hs.211612
3131	0.038612	putative translation initiation factor (SUI1), mRNA /cds=(148,489) /gb=NM_005801 /gi=20070210 /ug=Hs.150580 /len=1324	NM_005801	Hs.150580

3151	0.012673	hypothetical protein FLJ37440 (FLJ37440), mRNA /cds=(272,1591) /gb=NM_153214 /gi=23397470 /ug=Hs.355577 /len=2299	NM_153214	Hs.355577
3205	0.046607	hypothetical protein MGC2963 (MGC2963), mRNA /cds=(135,467) /gb=NM_031298 /gi=13775219 /ug=Hs.30011 /len=673	NM_031298	Hs.30011
3211	0.007929	chromobox 6 (CBX6), mRNA /cds=(30,1268) /gb=NM_014292 /gi=10140848 /ug=Hs.107374 /len=6014	NM_014292	Hs.107374
3212	0.046607	glutaminase (GLS), mRNA /cds=(231,2240) /gb=NM_014905 /gi=21361451 /ug=Hs.239189 /len=4799	NM_014905	Hs.239189
3229	0.038612	clone IMAGE:3457202, mRNA /gb=BC033700 /gi=22832845 /ug=Hs.324504 /len=2750	BC033700	Hs.324504
3230	0.005705	cDNA FLJ10004 fis, clone HEMBA1000076. /gb=AK000866 /gi=7021190 /ug=Hs.411490 /len=1974	AK000866	Hs.411490
3242	0.03394	KDEL (Lys-Asp-Glu-Leu) containing 1 (KDELC1), mRNA /cds=(338,1846) /gb=NM_024089 /gi=13129085 /ug=Hs.44970 /len=2082	NM_024089	Hs.44970
3248	0.017049	CDw92 antigen (CDW92), mRNA /cds=(43,2016) /gb=NM_080546 /gi=18034691 /ug=Hs.179902 /len=4301	NM_080546	Hs.179902
3312	0.038612	methyl-CpG binding domain protein 4 (MBD4), mRNA /cds=(177,1919) /gb=NM_003925 /gi=4505120 /ug=Hs.35947 /len=2470	NM_003925	Hs.35947
3348	0.012673	family with sequence similarity 8, member A1 (FAM8A1), mRNA /cds=(56,1297) /gb=NM_016255 /gi=7705267 /ug=Hs.95260 /len=4695	NM_016255	Hs.95260
3383	0.004048	mesenchymal stem cell protein DSC92 (NEUGRIN), mRNA /cds=(632,1291) /gb=NM_016645 /gi=7706195 /ug=Hs.323467 /len=1729	NM_016645	Hs.323467
3491	0.04114	cDNA: FLJ22071 fis, clone HEP11691. /gb=AK025724 /gi=10438333 /ug=Hs.422407 /len=2047	AK025724	Hs.422407
3749	0.046603	pericentrin 1 (PCNT1), mRNA /cds=(81,2051) /gb=NM_024844 /gi=13376258 /ug=Hs.184352 /len=2147	NM_024844	Hs.184352
3798	0.036213	chromodomain helicase DNA binding protein 4 (CHD4), mRNA /cds=(90,5828) /gb=NM_001273 /gi=4557452 /ug=Hs.74441 /len=6417	NM_001273	Hs.74441
4084	0.036213	hyperion gene, exons 1-50	AJ010770	
4113	0.036213	septin 10 mRNA, complete cds. /cds=(127,1680) /gb=AF146760 /gi=7688656 /ug=Hs.355455 /len=3018	AF146760	Hs.355455
4115	0.018325	WNT1 inducible signaling pathway protein 3 (WISP3), transcript variant 1, mRNA /cds=(111,1175) /gb=NM_003880 /gi=18491002 /ug=Hs.194678 /len=1307	NM_003880	Hs.194678
4133	0.049555	vimentin (VIM), mRNA /cds=(123,1523) /gb=NM_003380 /gi=4507894 /ug=Hs.297753 /len=1851	NM_003380	Hs.297753

4642	0.02782	mRNA; cDNA DKFZp686G1167 (from clone DKFZp686G1167) /gb=AL833600 /gi=21734246 /ug=Hs.7720 /len=8355	AL833600	Hs.7720
4646	0.043803	qh69a11.x1 Soares_fetal_liver_spleen_1NFLS_S1 cDNA clone IMAGE:1849916 3', mRNA sequence /clone=IMAGE:1849916 /clone_end=3' /gb=AI241467 /gi=3836864 /ug=Hs.303946 /len=398	AI241467	Hs.303946
4658	0.006737	HSPC125 protein (HSPC125), mRNA /cds=(80,607) /gb=NM_014165 /gi=7661785 /ug=Hs.5232 /len=725	NM_014165	Hs.5232
4699	0.049555	chromodomain protein, Y chromosome-like (CDYL), transcript variant 2, mRNA /cds=(336,1970) /gb=NM_170751 /gi=25777618 /ug=Hs.16081 /len=3474	NM_170751	Hs.16081
4728	0.013664	pp21 (LOC51186), mRNA /cds=(263,577) /gb=NM_016303 /gi=10047099 /ug=Hs.15984 /len=1038	NM_016303	Hs.15984
4729	0.013664	fer-1-like 3, myoferlin (C. elegans) (FER1L3), transcript variant 1, mRNA /cds=(89,6274) /gb=NM_013451 /gi=19718757 /ug=Hs.234680 /len=6829	NM_013451	Hs.234680
4741	0.019683	hypothetical protein MGC21981 (MGC21981), mRNA /cds=(66,764) /gb=NM_153267 /gi=23397567 /ug=Hs.131987 /len=1727	NM_153267	Hs.131987
4744	0.03394	seryl-tRNA synthetase (SARS), mRNA /cds=(76,1620) /gb=NM_006513 /gi=16306547 /ug=Hs.144063 /len=1942	NM_006513	Hs.144063
4764	0.025999	mRNA for KIAA1025 protein, partial cds. /cds=(1,5755) /gb=AB028948 /gi=20521733 /ug=Hs.4084 /len=8444	AB028948	Hs.4084
4801	0.036213	growth hormone inducible transmembrane protein (GHITM), mRNA /cds=(130,1089) /gb=NM_014394 /gi=7657479 /ug=Hs.433957 /len=2374	NM_014394	Hs.433957
4802	0.013664	A kinase (PRKA) anchor protein 1 (AKAP1), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA /cds=(63,1844) /gb=NM_139275 /gi=21493034 /ug=Hs.78921 /len=3801	NM_139275	Hs.78921
4807	0.029748	glypican 6 (GPC6), mRNA /cds=(616,2283) /gb=NM_005708 /gi=8051601 /ug=Hs.118407 /len=2760	NM_005708	Hs.118407
4815	0.043803	cDNA FLJ31247 fis, clone KIDNE2005296, weakly similar to ACTIN, CYTOPLASMIC 1. /gb=AK055809 /gi=16550632 /ug=Hs.170848 /len=2322	AK055809	Hs.170848
4821	0.012673	eukaryotic translation termination factor 1 (ETF1), mRNA /cds=(136,1449) /gb=NM_004730 /gi=4759033 /ug=Hs.77324 /len=3653	NM_004730	Hs.77324
4833	0.011744	ATP synthase, H transporting, mitochondrial F0 complex, subunit b, isoform 1 (ATP5F1), mRNA /cds=(98,868) /gb=NM_001688 /gi=21361564 /ug=Hs.81634 /len=1230	NM_001688	Hs.81634

4835	0.043803	tj81f06.x1 Soares_NSF_F8_9W_OT_PA_P_S1 cDNA clone IMAGE:2147939 3' similar to gb:M11050 GLUCOCORTICOID RECEPTOR, BETA mRNA sequence /clone=IMAGE:2147939 /clone_end=3' /gb=AI472273 /gi=4334363 /ug=Hs.228361 /len=555	AI472273	Hs.228361
4874	0.019683	ankyrin repeat domain 10 (ANKRD10), mRNA /cds=(136,1398) /gb=NM_017664 /gi=8923103 /ug=Hs.172572 /len=2509	NM_017664	Hs.172572
4880	0.025999	cDNA FLJ12776 fis, clone NT2RP2001678. /gb=AK022838 /gi=10434465 /ug=Hs.372558 /len=2629	AK022838	Hs.372558
4912	0.021125	ATP synthase, H transporting, mitochondrial F0 complex, subunit g (ATP5L), mRNA /cds=(60,371) /gb=NM_006476 /gi=21359881 /ug=Hs.107476 /len=580	NM_006476	Hs.107476
4913	0.021125	thymopoietin (TMPO) gene, exons 4 and 5, and complete cds for thymopoietin alpha	U18270	
4917	0.021125	mRNA for KIAA0257 gene, partial cds. /cds=(1,5419) /gb=D87446 /gi=1665780 /ug=Hs.75912 /len=6178	D87446	Hs.75912
4923	0.036213	KIAA1966 protein (KIAA1966), mRNA /cds=(492,2468) /gb=NM_133370 /gi=21166354 /ug=Hs.158184 /len=3248	NM_133370	Hs.158184
4995	0.03394	S100 calcium binding protein A10 (annexin II ligand, calpactin I, light polypeptide (p11)) (S100A10), mRNA /cds=(112,405) /gb=NM_002966 /gi=4506760 /ug=Hs.400250 /len=649	NM_002966	Hs.400250
5019	0.03394	desmin (DES), mRNA /cds=(81,1490) /gb=NM_001927 /gi=18105049 /ug=Hs.279604 /len=2236	NM_001927	Hs.279604
5059	0.031786	UDP-GalNAc:polypeptide N-acetylglactosaminyltransferase T10 (GalNAc-T10), mRNA /cds=(62,1735) /gb=NM_024572 /gi=13375743 /ug=Hs.15830 /len=2168	NM_024572	Hs.15830
5061	0.013664	wc09c01.x1 NCI_CGAP_Pr28 cDNA clone IMAGE:2314656 3' similar to gb:J05016 PROTEIN DISULFIDE ISOMERASE-RELATED PROTEIN PRECURSOR mRNA sequence /clone=IMAGE:2314656 /clone_end=3' /gb=AI674177 /gi=4874657 /ug=Hs.200089 /len=526	AI674177	Hs.200089
5066	0.022655	DKFZP586F1524 protein (DKFZP586F1524), mRNA /cds=(50,1156) /gb=NM_015584 /gi=7661671 /ug=Hs.241543 /len=2100	NM_015584	Hs.241543
5114	0.038612	hypothetical protein MGC11034 (MGC11034), mRNA /cds=(246,641) /gb=NM_031453 /gi=13899290 /ug=Hs.103378 /len=3301	NM_031453	Hs.103378
5134	0.010874	hypothetical protein DKFZp761B1514 (DKFZp761B1514), mRNA /cds=(73,1029) /gb=NM_032288 /gi=14150032 /ug=Hs.177537 /len=3453	NM_032288	Hs.177537

5142	0.04114	solute carrier family 29 (nucleoside transporters), member 1 (SLC29A1), mRNA /cds=(179,1549) /gb=NM_004955 /gi=4826715 /ug=Hs.25450 /len=2162	NM_004955	Hs.25450
5143	0.049555	excision repair cross-complementing rodent repair deficiency, complementation group 3 (xeroderma pigmentosum group B complementing) (ERCC3), mRNA /cds=(96,2444) /gb=NM_000122 /gi=4557562 /ug=Hs.77929 /len=2751	NM_000122	Hs.77929
5145	0.043803	UI-E-C10-aah-e-03-0-UI.r1 UI-E-C10 cDNA clone UI-E-C10-aah-e-03-0-UI 5', mRNA sequence /clone=UI-E-C10-aah-e-03-0-UI /clone_end=5' /gb=BM690376 /gi=19003634 /ug=Hs.284207 /len=1151	BM690376	Hs.284207
5147	0.022655	blue cone opsin gene, complete cds	L32835	
5157	0.049555	Kallmann syndrome 1 sequence (KAL1), mRNA /cds=(151,2193) /gb=NM_000216 /gi=4557682 /ug=Hs.89591 /len=6314	NM_000216	Hs.89591
5221	0.043803	ATP synthase, H transporting, mitochondrial F0 complex, subunit b, isoform 1 (ATP5F1), mRNA /cds=(98,868) /gb=NM_001688 /gi=21361564 /ug=Hs.81634 /len=1230	NM_001688	Hs.81634
5293	0.025999	chromosome 6 open reading frame 48 (C6orf48), mRNA /cds=(42,422) /gb=NM_016947 /gi=8393383 /ug=Hs.109798 /len=711	NM_016947	Hs.109798
5325	0.031786	eukaryotic translation elongation factor 2 (EEF2), mRNA /cds=(69,2645) /gb=NM_001961 /gi=25453476 /ug=Hs.75309 /len=3148	NM_001961	Hs.75309
5365	0.036213	cDNA FLJ31667 fis, clone NT2RI2004840. /gb=AK056229 /gi=16551572 /ug=Hs.48692 /len=2052	AK056229	Hs.48692
5367	0.049555	hypothetical protein MGC4415 (MGC4415), mRNA /cds=(154,675) /gb=NM_031484 /gi=13899343 /ug=Hs.209614 /len=3243	NM_031484	Hs.209614
5458	0.011744	mRNA for KIAA0993 protein, partial cds	AB023210	Hs.198135
5476	0.029748	meningioma expressed antigen 6 (coiled-coil proline-rich) (MGEA6), mRNA /cds=(315,2729) /gb=NM_005930 /gi=5174560 /ug=Hs.117242 /len=3676	NM_005930	Hs.117242
5495	0.036213	clone IMAGE:4798349, mRNA /gb=BC045794 /gi=28277189 /ug=Hs.29464 /len=2717	BC045794	Hs.29464
5505	0.02782	SECIS binding protein 2 (SBP2), mRNA /cds=(58,2622) /gb=NM_024077 /gi=21359954 /ug=Hs.288141 /len=3457	NM_024077	Hs.288141
5515	0.046607	cell cycle progression 8 protein (CPR8), mRNA /cds=(13,1140) /gb=NM_004748 /gi=4758047 /ug=Hs.82506 /len=1856	NM_004748	Hs.82506
5551	0.043803	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit (HADHA), mRNA /cds=(35,2326) /gb=NM_000182 /gi=20127407 /ug=Hs.75860 /len=2972	NM_000182	Hs.75860

5634	0.015849	yc38b10.s1 Stratagene liver (#937224) cDNA clone IMAGE:82939 3' similar to gb:M24398 parathymosin mRNA, complete cds. mRNA sequence /clone=IMAGE:82939 /clone_end=3' /gb=T69405 /gi=680553 /ug=Hs.427073 /len=465	T69405	Hs.427073
5707	0.017049	nucleolar autoantigen (55kD) similar to rat synaptonemal complex protein (SC65), mRNA /cds=(12,1325) /gb=NM_006455 /gi=5454037 /ug=Hs.446459 /len=2347	NM_006455	Hs.446459
5782	0.029748	DNA sequence from clone RP11-305F14 on chromosome 9 Contains part of a novel gene for astrotactin 2, ESTs, STSs, GSSs and a CpG island, complete sequence	AL157829	
5824	0.038612	activated RNA polymerase II transcription cofactor 4 (PC4), mRNA /cds=(57,440) /gb=NM_006713 /gi=19923783 /ug=Hs.349506 /len=1336	NM_006713	Hs.349506
5843	0.024278	MYE4197a Myeloma (MYE) cDNA library cDNA, mRNA sequence /gb=BF174993 /gi=13441207 /ug=Hs.332023 /len=338	BF174993	Hs.332023
5855	0.043803	coactosin-like 1 (Dictyostelium) (COTL1), mRNA /cds=(150,578) /gb=NM_021149 /gi=23510452 /ug=Hs.289092 /len=1850	NM_021149	Hs.289092
5861	0.021123	mRNA for KIAA0338 gene, partial cds. /cds=(1,2807) /gb=AB002336 /gi=2224616 /ug=Hs.26395 /len=6263	AB002336	Hs.26395
5874	0.03394	BX104663 Soares_pregnant_uterus_NbHPU cDNA clone IMAGp998J071170, mRNA sequence /clone=IMAGp998J071170;_IMAGE:491622 /gb=BX104663 /gi=27846238 /ug=Hs.426388 /len=748	BX104663	Hs.426388
5917	0.025999	no match		
5924	0.046607	B-cell receptor-associated protein BAP29 (BAP29), mRNA /cds=(47,775) /gb=NM_018844 /gi=9994198 /ug=Hs.27135 /len=1085	NM_018844	Hs.27135
5960	0.024278	zinc finger protein ANC_2H01 (LOC51193), mRNA /cds=(446,1903) /gb=NM_016331 /gi=7705934 /ug=Hs.22879 /len=3013	NM_016331	Hs.22879
5989	0.043803	CDA02 protein (CDA02), mRNA /cds=(3,1832) /gb=NM_032025 /gi=14042940 /ug=Hs.332404 /len=2179	NM_032025	Hs.332404
5996	0.031786	mitochondrial ribosomal protein L40 (MRPL40), nuclear gene encoding mitochondrial protein, mRNA /cds=(43,663) /gb=NM_003776 /gi=26638658 /ug=Hs.431307 /len=787	NM_003776	Hs.431307
6002	0.029748	proteasome (prosome, macropain) activator subunit 2 (PA28 beta) (PSME2), mRNA /cds=(66,785) /gb=NM_002818 /gi=4506236 /ug=Hs.433810 /len=828	NM_002818	Hs.433810
6088	0.005243	cyclin G associated kinase (GAK), mRNA /cds=(1,3936) /gb=NM_005255 /gi=4885250 /ug=Hs.153227 /len=4331	NM_005255	Hs.153227
6135	0.021125	cyclin-dependent kinase 4 (CDK4), transcript variant 1, mRNA /cds=(228,1139) /gb=NM_000075 /gi=16936531 /ug=Hs.95577 /len=1474	NM_000075	Hs.95577

6228	0.007312	prefoldin 2 (PFDN2), mRNA /cds=(31,495) /gb=NM_012394 /gi=12408674 /ug=Hs.298229 /len=644	NM_012394	Hs.298229
6353	0.029748	clone IMAGE:3681106, mRNA, partial cds /cds=(1,71) /gb=BC014579 /gi=15779008 /ug=Hs.348647 /len=1232	BC014579	Hs.348647
6471	0.024278	hydroxysteroid (11-beta) dehydrogenase 1 (HSD11B1), mRNA /cds=(95,973) /gb=NM_005525 /gi=5031764 /ug=Hs.275215 /len=1375	NM_005525	Hs.275215
6550	0.03394	t-complex-associated-testis-expressed 1-like 1 (TCTEL1), mRNA /cds=(1,342) /gb=NM_006519 /gi=5730084 /ug=Hs.266940 /len=713	NM_006519	Hs.266940
6568	0.029748	hypothetical protein HSPC132 (HSPC132), mRNA /cds=(4,234) /gb=NM_016399 /gi=7705466 /ug=Hs.69499 /len=1171	NM_016399	Hs.69499
6599	0.024278	N-myc downstream regulated gene 1 (NDRG1), mRNA /cds=(111,1295) /gb=NM_006096 /gi=5174656 /ug=Hs.75789 /len=3020	NM_006096	Hs.75789
6610	0.036213	Similar to RIKEN cDNA 3830613O22 gene, clone IMAGE:5551209, mRNA, partial cds /cds=(282,4079) /gb=BC035645 /gi=23272851 /ug=Hs.356876 /len=4079	BC035645	Hs.356876
6613	0.04114	major histocompatibility complex, class II, DR beta 1 (HLA-DRB1), mRNA /cds=(63,863) /gb=NM_002124 /gi=4504410 /ug=Hs.375570 /len=1182	NM_002124	Hs.375570
6614	0.026894	actin related protein 2/3 complex, subunit 2, 34kDa (ARPC2), transcript variant 1, mRNA /cds=(113,1015) /gb=NM_152862 /gi=23238210 /ug=Hs.83583 /len=1462	NM_152862	Hs.83583
6636	0.025999	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 15 (DDX15), mRNA /cds=(162,2603) /gb=NM_001358 /gi=4557516 /ug=Hs.5683 /len=3028	NM_001358	Hs.5683
6672	0.013664	polymerase (RNA) II (DNA directed) polypeptide A, 220kDa (POLR2A), mRNA /cds=(387,6299) /gb=NM_000937 /gi=14589948 /ug=Hs.171880 /len=6732	NM_000937	Hs.171880
6677	0.018325	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 5 (SLC25A5), nuclear gene encoding mitochondrial protein, mRNA /cds=(69,965) /gb=NM_001152 /gi=4502098 /ug=Hs.79172 /len=1225	NM_001152	Hs.79172
6679	0.04114	hypothetical protein PP1057 (PP1057), mRNA /cds=(498,971) /gb=NM_031285 /gi=13775195 /ug=Hs.108557 /len=2030	NM_031285	Hs.108557
6699	0.021125	CGI-26 protein (CGI-26), mRNA /cds=(20,1030) /gb=NM_015954 /gi=7705717 /ug=Hs.24332 /len=1561	NM_015954	Hs.24332
6701	0.049555	inositol 1,4,5-triphosphate receptor, type 2 (ITPR2), mRNA	NM_002223	

6717	0.015849	fer-1-like 3, myoferlin (<i>C. elegans</i>) (FER1L3), transcript variant 1, mRNA /cds=(89,6274) /gb=NM_013451 /gi=19718757 /ug=Hs.234680 /len=6829	NM_013451	Hs.234680
6750	0.029748	ADP-ribosylation-like factor 6 interacting protein 4 (ARL6IP4), mRNA /cds=(63,719) /gb=NM_016638 /gi=7706183 /ug=Hs.103561 /len=952	NM_016638	Hs.103561
6753	0.031786	mRNA; cDNA DKFZp434A012 (from clone DKFZp434A012) /gb=AL096752 /gi=5419888 /ug=Hs.306327 /len=2248	AL096752	Hs.306327
6767	0.021125	ATP synthase, H transporting, mitochondrial F0 complex, subunit g (ATP5L), mRNA /cds=(60,371) /gb=NM_006476 /gi=21359881 /ug=Hs.107476 /len=580	NM_006476	Hs.107476
6794	0.046607	chromosome 13 open reading frame 12 (C13orf12), mRNA /cds=(76,501) /gb=NM_015932 /gi=21361533 /ug=Hs.279813 /len=1352	NM_015932	Hs.279813
6857	0.025999	GRB2-associated binding protein 1 (GAB1), mRNA /cds=(122,2206) /gb=NM_002039 /gi=4503850 /ug=Hs.239706 /len=2467	NM_002039	Hs.239706
6885	0.006202	CDK2-associated protein 1 (CDK2AP1), mRNA /cds=(523,870) /gb=NM_004642 /gi=17978492 /ug=Hs.433201 /len=1627	NM_004642	Hs.433201
6900	0.004417	developmentally regulated GTP binding protein 1 (DRG1), mRNA /cds=(66,1169) /gb=NM_004147 /gi=4758795 /ug=Hs.115242 /len=1383	NM_004147	Hs.115242
6931	0.010874	Mov10, Moloney leukemia virus 10, (mouse) (MOV10), mRNA /cds=(71,3082) /gb=NM_020963 /gi=14211539 /ug=Hs.20725 /len=3328	NM_020963	Hs.20725
6933	0.041114	mRNA full length insert cDNA clone EUROIMAGE 2004632	AL389975	Hs.28219
6957	0.021125	ATP synthase, H transporting, mitochondrial F1 complex, O subunit (oligomycin sensitivity conferring protein) (ATP5O), mRNA /cds=(37,678) /gb=NM_001697 /gi=4502302 /ug=Hs.433960 /len=772	NM_001697	Hs.433960
6964	0.012673	vacuolar protein sorting 4A (yeast) (VPS4A), mRNA /cds=(129,1442) /gb=NM_013245 /gi=17865806 /ug=Hs.234839 /len=2211	NM_013245	Hs.234839
7045	0.038612	methionine adenosyltransferase II, beta (MAT2B), mRNA /cds=(73,1077) /gb=NM_013283 /gi=20127525 /ug=Hs.54642 /len=2054	NM_013283	Hs.54642
7124	0.041114	annexin A5 (ANXA5), mRNA /cds=(193,1155) /gb=NM_001154 /gi=4809273 /ug=Hs.300711 /len=1630	NM_001154	Hs.300711
7149	0.043803	solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 3 (SLC25A3), nuclear gene encoding mitochondrial protein, transcript variant 1b, mRNA /cds=(49,1134) /gb=NM_002635 /gi=4505774 /ug=Hs.78713 /len=1330	NM_002635	Hs.78713

7487	0.04114	protease, serine, 11 (IGF binding) (PRSS11), mRNA /cds=(49,1491) /gb=NM_002775 /gi=21327712 /ug=Hs.75111 /len=2039	NM_002775	Hs.75111
7510	0.018325	hypothetical protein MGC14327 (MGC14327), mRNA /cds=(225,635) /gb=NM_053045 /gi=16596685 /ug=Hs.231029 /len=1576	NM_053045	Hs.231029
7543	0.038612	duplicated clone on array		
7619	0.022655	mannosidase, alpha, class 1A, member 1 (MAN1A1), mRNA /cds=(443,2404) /gb=NM_005907 /gi=24497518 /ug=Hs.25253 /len=4139	NM_005907	Hs.25253
7646	0.004814	postsynaptic protein CRIPT (CRIPT), mRNA /cds=(88,393) /gb=NM_014171 /gi=7661797 /ug=Hs.39733 /len=1155	NM_014171	Hs.39733
7662	0.008591	core promoter element binding protein (COPEB), mRNA /cds=(118,969) /gb=NM_001300 /gi=9961346 /ug=Hs.285313 /len=1470	NM_001300	Hs.285313
7671	0.025999	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA /cds=(266,2413) /gb=NM_004728 /gi=13787208 /ug=Hs.169531 /len=3319	NM_004728	Hs.169531
7725	0.046607	proteasome (prosome, macropain) subunit, beta type, 2 (PSMB2), mRNA /cds=(111,716) /gb=NM_002794 /gi=22538463 /ug=Hs.432607 /len=850	NM_002794	Hs.432607
7727	0.006737	hypothetical protein FLJ13081 (FLJ13081), mRNA /cds=(171,2099) /gb=NM_024834 /gi=13376242 /ug=Hs.180638 /len=4113	NM_024834	Hs.180638
7728	0.017049	Down syndrome critical region gene 5 (DSCR5), transcript variant 3, mRNA /cds=(342,668) /gb=NM_016430 /gi=24497594 /ug=Hs.408790 /len=875	NM_016430	Hs.408790
7759	0.038612	hypothetical protein FLJ14825 (FLJ14825), mRNA /cds=(32,1174) /gb=NM_032847 /gi=14249567 /ug=Hs.334824 /len=1286	NM_032847	Hs.334824
7892	0.022655	BAC clone RP11-36B15 from 4, complete sequence	AC027607	
7897	0.018325	mRNA for FLJ00245 protein	AK074172	Hs.244343
8049	0.022655	KIAA0073 protein (KIAA0073), mRNA /cds=(4,1944) /gb=NM_015342 /gi=24308048 /ug=Hs.1191 /len=2110	NM_015342	Hs.1191
8058	0.04114	farnesyltransferase, CAAX box, alpha (FNTA), mRNA /cds=(7,1146) /gb=NM_002027 /gi=4503770 /ug=Hs.356463 /len=1644	NM_002027	Hs.356463
8087	0.036213	cDNA FLJ36527 fis, clone TRACH2003941. /gb=AK093846 /gi=21752790 /ug=Hs.378776 /len=2526	AK093846	Hs.378776
8089	0.017049	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4 (B4GALT4), mRNA /cds=(311,1345) /gb=NM_003778 /gi=13929469 /ug=Hs.13225 /len=2167	NM_003778	Hs.13225
8294	0.043803	DNA sequence from clone RP11-21M8 on chromosome 10, complete sequence	AL157711	
8306	0.029748	hypothetical protein FLJ20313 (FLJ20313), mRNA /cds=(345,1700) /gb=NM_017762 /gi=8923296 /ug=Hs.126721 /len=2226	NM_017762	Hs.126721

8403	0.02782	chromosome 18, clone RP11-7L24, complete sequence	AC013587	
8419	0.036213	mRNA; cDNA DKFZp586B1922 (from clone DKFZp586B1922) /gb=AL049450 /gi=4500236 /ug=Hs.184779 /len=1433	AL049450	Hs.184779
8434	0.043803	wq37a10.x1 NCI_CGAP_GC6 cDNA clone IMAGE:2473434 3' similar to contains element MER18 repetitive element ;, mRNA sequence /clone=IMAGE:2473434 /clone_end=3' /gb=AI948831 /gi=5741141 /ug=Hs.224268 /len=430	AI948831	Hs.224268
8571	0.017049	AGENCOURT_6758988 NIH_MGC_115 cDNA clone IMAGE:5755234 5', mRNA sequence /clone=IMAGE:5755234 /clone_end=5' /gb=BQ067651 /gi=19896697 /ug=Hs.204354 /len=1385	BQ067651	Hs.204354
8580	0.029748	BX096173 Soares_testis_NHT cDNA clone IMAGp998F151793, mRNA sequence /clone=IMAGp998F151793 ;_IMAGE:730766 /gb=BX096173 /gi=27842669 /ug=Hs.188780 /len=556	BX096173	Hs.188780
8718	0.018325	im93f12.x1 insulinoma cDNA clone IMAGE:6122639 3', mRNA sequence /clone=IMAGE:6122639 /clone_end=3' /gb=BU786820 /gi=23833860 /ug=Hs.145958 /len=625	BU786820	Hs.145958
8869	0.022655	no match		
8879	0.007929	no match		
8925	0.03394	DNA sequence from clone RP11-550P23 on chromosome 13, complete sequence	AL353648	
8928	0.015849	cDNA: FLJ23313 fis, clone HEP11919. /gb=AK026966 /gi=10439954 /ug=Hs.10862 /len=2527	AK026966	Hs.10862
8978	0.022655	Similar to procollagen, type V, alpha 2, clone IMAGE:3613441, mRNA /gb=BC014149 /gi=15559579 /ug=Hs.162411 /len=1335	BC014149	Hs.162411
9050	0.009301	cDNA FLJ25252 fis, clone STM03814. /gb=AK057981 /gi=16553973 /ug=Hs.16979 /len=2005	AK057981	Hs.16979
9108	0.018325	UI-H-EI0-aye-c-17-0-UI.s1 NCI_CGAP_EI0 cDNA clone UI-H-EI0-aye-c-17-0-UI 3', mRNA sequence /clone=UI-H-EI0-aye-c-17-0-UI /clone_end=3' /gb=CA447385 /gi=24811805 /ug=Hs.420740 /len=812	CA447385	Hs.420740
9185	0.019683	UI-1-BB1p-akd-c-08-0-UI.s1 NCI_CGAP_PI6 cDNA clone UI-1-BB1p-akd-c-08-0-UI 3', mRNA sequence /clone=UI-1-BB1p-akd-c-08-0-UI /clone_end=3' /gb=BQ026195 /gi=19761474 /ug=Hs.308520 /len=1138	BQ026195	Hs.308520
9200	0.031786	602387746F1 NIH_MGC_93 cDNA clone IMAGE:4516739 5', mRNA sequence /clone=IMAGE:4516739 /clone_end=5' /gb=BG287971 /gi=13042340 /ug=Hs.303110 /len=749	BG287971	Hs.303110
9406	0.038612	FK506 binding protein 9, 63 kDa (FKBP9), mRNA /cds=(457,885) /gb=NM_007270 /gi=24307926 /ug=Hs.302749 /len=2517	NM_007270	Hs.302749

9426	0.012673	clone IMAGE:4130494, mRNA /gb=BC023543 /gi=23270740 /ug=Hs.112844 /len=4567	BC023543	Hs.112844
9451	0.019683	phospholipase A2-activating protein (PLAA), mRNA /cds=(29,2245) /gb=NM_004253 /gi=21361288 /ug=Hs.27182 /len=3240	NM_004253	Hs.27182
9558	0.036213	Gene 33/Mig-6 (MIG-6), mRNA /cds=(213,1601) /gb=NM_018948 /gi=21314673 /ug=Hs.11169 /len=3099	NM_018948	Hs.11169
9561	0.010061	phosphatidylinositol transfer protein, beta (PITPNB), mRNA /cds=(40,855) /gb=NM_012399 /gi=19923401 /ug=Hs.7370 /len=2906	NM_012399	Hs.7370
9633	0.046607	U7 snRNP-specific Sm-like protein LSM10 (LSM10), mRNA /cds=(151,522) /gb=NM_032881 /gi=14249631 /ug=Hs.3496 /len=869	NM_032881	Hs.3496
10066	0.043803	CGI-32 protein (CGI-32), mRNA /cds=(103,924) /gb=NM_015960 /gi=7705727 /ug=Hs.16606 /len=1323	NM_015960	Hs.16606
10382	0.014722	hypothetical protein DKFZp761N0624 (DKFZp761N0624), mRNA /cds=(113,1444) /gb=NM_032295 /gi=14150046 /ug=Hs.21893 /len=2973	NM_032295	Hs.21893
10472	0.029748	mRNA; cDNA DKFZp434L0823 (from clone DKFZp434L0823) /gb=AL833965 /gi=21739550 /ug=Hs.440644 /len=2628	AL833965	Hs.440644
10485	0.043803	cDNA FLJ36429 fis, clone THYMU2011573. /gb=AK093748 /gi=21752675 /ug=Hs.378821 /len=1901	AK093748	Hs.378821
10593	0.046607	twisted gastrulation 1 (Drosophila) (TWSG1), mRNA /cds=(106,777) /gb=NM_020648 /gi=21314788 /ug=Hs.247302 /len=3693	NM_020648	Hs.247302
10621	0.036213	chromosome 5 clone RP11-147O19, complete sequence	AC093250	
10729	0.022655	myc-induced nuclear antigen, 53 kDa (MINA53), transcript variant 2, mRNA /cds=(214,1608) /gb=NM_032778 /gi=23346417 /ug=Hs.23294 /len=2221	NM_032778	Hs.23294
10771	0.03394	HUM515D03B Clontech placenta polyA mRNA (#6518) cDNA clone GEN-515D03 5', mRNA sequence /clone=GEN-515D03 /clone_end=5' /gb=D58893 /gi=968527 /ug=Hs.335953 /len=365	D58893	Hs.335953
10845	0.015849	spinocerebellar ataxia 2 (olivopontocerebellar ataxia 2, autosomal dominant, ataxin 2) (SCA2), mRNA /cds=(163,4101) /gb=NM_002973 /gi=4506794 /ug=Hs.76253 /len=4481	NM_002973	Hs.76253
10908	0.03394	chromosome 5 clone CTB-113P19, complete sequence	AC011374	
10916	0.02782	cDNA FLJ34008 fis, clone FCBBF1000425	AK091327	Hs.356216
10978	0.014722	cDNA FLJ31827 fis, clone NT2RP6000100, moderately similar to ZINC FINGER PROTEIN 41. /cds=(474,1694) /gb=AK056389 /gi=16551782 /ug=Hs.378531 /len=3180	AK056389	Hs.378531

11050	0.046607	clone IMAGE:4693260, mRNA /gb=BC017972 /gi=17389932 /ug=Hs.24444 /len=1637	BC017972	Hs.24444
11066	0.03394	mRNA; cDNA DKFZp667D083 (from clone DKFZp667D083)	AL713688	
11069	0.046607	3 RP11-159G9 (Roswell Park Cancer Institute BAC Library) complete sequence	AC119733	
11104	0.013664	clone IMAGE:5268031, mRNA /gb=BC040578 /gi=26251832 /ug=Hs.287864 /len=3284	BC040578	Hs.287864
11204	0.010874	BAC clone RP11-489G11 from 4, complete sequence	AC093849	
11241	0.025999	cDNA FLJ32072 fis, clone OCBBF1000130	AK056634	
11320	0.049555	nuclear fragile X mental retardation protein interacting protein 1 (NUFIP1), mRNA /cds=(1,1488) /gb=NM_012345 /gi=6912541 /ug=Hs.120247 /len=3463	NM_012345	Hs.120247
11329	0.010061	similar to CG9578 gene product (MGC3794), mRNA /cds=(146,964) /gb=NM_152902 /gi=23097249 /ug=Hs.137576 /len=1314	NM_152902	Hs.137576
11358	0.046607	v-raf-1 murine leukemia viral oncogene 1 (RAF1), mRNA /cds=(130,2076) /gb=NM_002880 /gi=4506400 /ug=Hs.349650 /len=2977	NM_002880	Hs.349650
11448	0.018325	zinc finger protein 23 (KOX 16) (ZNF23), mRNA /cds=(815,2746) /gb=NM_145911 /gi=23308736 /ug=Hs.376810 /len=3271	NM_145911	Hs.376810
11492	0.046607	chromosome 3, olfactory receptor pseudogene cluster 1, complete sequence, and myosin light chain kinase (MLCK) pseudogene, partial sequence	AF042089	
11631	0.046607	chromosome 11, clone RP11-449L13, complete sequence	AC100865	
11693	0.006737	cell adhesion molecule-related/down-regulated by oncogenes (CDON), mRNA /cds=(1,3723) /gb=NM_016952 /gi=8393083 /ug=Hs.159565 /len=3986	NM_016952	Hs.159565
11744	0.04114	mitofusin 1 (MFN1), transcript variant 1, mRNA /cds=(84,2309) /gb=NM_033540 /gi=16117784 /ug=Hs.197877 /len=3275	NM_033540	Hs.197877
11769	0.015849	golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA /cds=(132,2327) /gb=NM_005113 /gi=4826747 /ug=Hs.241572 /len=2838	NM_005113	Hs.241572
11772	0.02782	legumain (LGMN), mRNA /cds=(142,1443) /gb=NM_005606 /gi=21914880 /ug=Hs.18069 /len=1981	NM_005606	Hs.18069
11811	0.003707	cDNA FLJ11481 fis, clone HEMBA1001803. /gb=AK021543 /gi=10432744 /ug=Hs.135159 /len=1539	AK021543	Hs.135159
11915	0.046607	paraoxonase 2 (PON2), mRNA /cds=(33,1097) /gb=NM_000305 /gi=4505952 /ug=Hs.169857 /len=1600	NM_000305	Hs.169857
11937	0.013664	Mus musculus dual specificity phosphatase 14 (Dusp14), mRNA	XM_282991	
11970	0.036213	mRNA for KIAA1728 protein, partial cds. /cds=(1,4937) /gb=AB051515 /gi=12698000 /ug=Hs.252748 /len=6585	AB051515	Hs.252748

12114	0.024278	nac89f07.x1 NCI_CGAP_Brn23 cDNA clone IMAGE:3441564 3', mRNA sequence /clone=IMAGE:3441564 /clone_end=3' /gb=BG054792 /gi=12511863 /ug=Hs.157695 /len=581	BG054792	Hs.157695
12143	0.015849	pVHL-interacting deubiquitinating enzyme 1 (VDU1), mRNA /cds=(262,2997) /gb=NM_015017 /gi=21489974 /ug=Hs.173694 /len=4323	NM_015017	Hs.173694
12197	0.025999	DKFZp586E2017_r1 586 (synonym: hute1) cDNA clone DKFZp586E2017 5', mRNA sequence /clone=DKFZp586E2017 /clone_end=5' /gb=AL046885 /gi=5936275 /ug=Hs.413463 /len=640	AL046885	Hs.413463
12243	0.007929	mRNA; cDNA DKFZp313P0434 (from clone DKFZp313P0434) /gb=AL832702 /gi=21733281 /ug=Hs.125019 /len=2995	AL832702	Hs.125019
12441	0.011744	cDNA FLJ12411 fis, clone MAMMA1002964	AK022473	
12514	0.029748	DCBCQH10 DCB cDNA, mRNA sequence /gb=BU198777 /gi=22717083 /ug=Hs.50273 /len=867	BU198777	Hs.50273
12517	0.019683	UI-H-DP0-avt-a-17-0-UI.s1 NCI_CGAP_Fs1 cDNA clone IMAGE:5883928 3', mRNA sequence /clone=IMAGE:5883928 /clone_end=3' /gb=BQ000272 /gi=19725172 /ug=Hs.371473 /len=1051	BQ000272	Hs.371473
12538	0.046607	chromosome 8, clone CTD-3046C4, complete sequence	AC103688	
12609	0.031786	Similar to hypothetical protein FLJ31322, clone IMAGE:5296647, mRNA /gb=BC045189 /gi=28277118 /ug=Hs.350001 /len=2971	BC045189	Hs.350001
12629	0.02782	no match		
12727	0.012673	BAC clone RP11-745K4 from 4, complete sequence	AC097108	
12756	0.011744	DNA sequence from clone RP11-391H12 on chromosome 13, complete sequence	AL136221	
12759	0.006202	UI-H-BI1-aee-g-08-0-UI.s1 NCI_CGAP_Sub3 cDNA clone IMAGE:2719286 3', mRNA sequence /clone=IMAGE:2719286 /clone_end=3' /gb=AW139957 /gi=6144675 /ug=Hs.396536 /len=744	AW139957	Hs.396536
12819	0.024278	BAC clone RP11-84C2 from 2, complete sequence	AC073415	
12822	0.038612	yu41a04.y5 Soares ovary tumor NbHOT cDNA clone IMAGE:236334 5' similar to contains Alu repetitive element; mRNA sequence /clone=IMAGE:236334 /clone_end=5' /gb=AI820819 /gi=5439898 /ug=Hs.193116 /len=567	AI820819	Hs.193116
12852	0.038612	oy68d07.x1 NCI_CGAP_CLL1 cDNA clone IMAGE:1670989 3', mRNA sequence /clone=IMAGE:1670989 /clone_end=3' /gb=AI085586 /gi=3424009 /ug=Hs.276342 /len=357	AI085586	Hs.276342
12896	0.029748	cDNA FLJ30298 fis, clone BRACE2003172. /gb=AK054860 /gi=16549479 /ug=Hs.351546 /len=2659	AK054860	Hs.351546
12906	0.049555	chromosome 10 clone RP11-346D6, complete sequence	AC009986	

12967	0.014722	to16a05.x1 NCI_CGAP_Ut2 cDNA clone IMAGE:2179184 3', mRNA sequence /clone=IMAGE:2179184 /clone_end=3' /gb=AI565553 /gi=4524010 /ug=Hs.309582 /len=404	AI565553	Hs.309582
12968	0.04114	cDNA FLJ37403 fis, clone BRAMY2028183. /gb=AK094722 /gi=21753837 /ug=Hs.117747 /len=2781	AK094722	Hs.117747
13068	0.018325	Alien PCR 4		
13114	0.049555	eukaryotic translation initiation factor 2C, 1 (EIF2C1), mRNA /cds=(214,2787) /gb=NM_012199 /gi=6912351 /ug=Hs.14520 /len=7478	NM_012199	Hs.14520
13175	0.049555	chromosome 1 open reading frame 9 (C1orf9), mRNA /cds=(125,4342) /gb=NM_016227 /gi=7705321 /ug=Hs.108636 /len=5919	NM_016227	Hs.108636
13193	0.031786	protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform (PPP2CA), mRNA /cds=(210,1139) /gb=NM_002715 /gi=4506016 /ug=Hs.91773 /len=2181	NM_002715	Hs.91773
13246	0.019683	KIAA1309 protein (KIAA1309), mRNA /cds=(211,2025) /gb=NM_033495 /gi=15741229 /ug=Hs.348262 /len=3119	NM_033495	Hs.348262
13277	0.015847	mRNA for KIAA1134 protein, partial cds. /cds=(1,1838) /gb=AB032960 /gi=6382011 /ug=Hs.18282 /len=5148	AB032960	Hs.18282
13488	0.046607	hypothetical protein MGC12262 (MGC12262), mRNA /cds=(566,784) /gb=NM_032696 /gi=14249285 /ug=Hs.334672 /len=868	NM_032696	Hs.334672
13508	0.036213	wf97f03.x1 Soares_NSF_F8_9W_OT_PA_P_S1 cDNA clone IMAGE:2363549 3', mRNA sequence /clone=IMAGE:2363549 /clone_end=3' /gb=AI808816 /gi=5395382 /ug=Hs.270591 /len=552	AI808816	Hs.270591
13511	0.036213	partial GLOMULIN gene for glomulin, exons 2-3 and join CDS	AJ302727	
13749	0.012673	SH3-domain GRB2-like endophilin B1 (SH3GLB1), mRNA /cds=(82,1179) /gb=NM_016009 /gi=21359904 /ug=Hs.136309 /len=1561	NM_016009	Hs.136309
13824	0.003392	Protein P3 (P3), mRNA /cds=(493,1926) /gb=NM_019848 /gi=10938005 /ug=Hs.72980 /len=2123	NM_019848	Hs.72980
13858	0.018325	mRNA; cDNA DKFZp761L03121 (from clone DKFZp761L03121) /cds=(1,3991) /gb=AL833838 /gi=21739301 /ug=Hs.7316 /len=4583	AL833838	Hs.7316
13987	0.04114	mRNA; cDNA DKFZp761C169 (from clone DKFZp761C169); partial cds /cds=(997,2475) /gb=AL161991 /gi=7328122 /ug=Hs.71252 /len=3324	AL161991	Hs.71252
14246	0.031786	mRNA; cDNA DKFZp564C2063 (from clone DKFZp564C2063) /gb=AL117595 /gi=5912159 /ug=Hs.4055 /len=1444	AL117595	Hs.4055
14249	0.036213	mitochondrion, complete genome	NC_001807	

14343	0.043803	UI-H-FH1-bfp-m-06-0-UI.s1 NCI_CGAP_FH1 cDNA clone UI-H-FH1-bfp-m-06-0-UI 3', mRNA sequence /clone=UI-H-FH1-bfp-m-06-0-UI /clone_end=3' /gb=BU619573 /gi=23285788 /ug=Hs.312629 /len=1168	BU619573	Hs.312629
14416	0.038612	chromosome 17, clone CTD-2026D20, complete sequence	AC040934	
14462	0.025999	chromosome 3 clone RP11-11L6, complete sequence	AC093412	
14529	0.046607	hypothetical protein FLJ23033 (FLJ23033), mRNA /cds=(108,1826) /gb=NM_024686 /gi=13375957 /ug=Hs.96423 /len=2115	NM_024686	Hs.96423
14614	0.043803	EST(yq95a02.r1 Soares fetal liver spleen 1NFLS cDNA clone IMAGE:203498 5' similar to contains Alu repetitive element)	H56096.1	
14629	0.029748	UI-H-ED1-axs-i-05-0-UI.s1 NCI_CGAP_ED1 cDNA clone IMAGE:5833036 3', mRNA sequence /clone=IMAGE:5833036 /clone_end=3' /gb=BQ014114 /gi=19739015 /ug=Hs.195045 /len=1024	BQ014114	Hs.195045
14631	0.014722	DNA sequence from clone RP11-434B7 on chromosome 1, complete sequence	AL583826	
14754	0.02782	mRNA; cDNA DKFZp313P0434 (from clone DKFZp313P0434) /gb=AL832702 /gi=21733281 /ug=Hs.125019 /len=2995	AL832702	Hs.125019
14856	0.049555	mRNA; cDNA DKFZp434K087 (from clone DKFZp434K087)	AL133031	Hs.193602